

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 09/889,331B
Source: 1FW16
Date Processed by STIC: 1/14/05

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 01/14/2005

PATENT APPLICATION: US/09/889,331B

TIME: 17:11:43

Input Set : A:\249-167 subst seq listing.txt

Output Set: N:\CRF4\01142005\I889331B.raw

3 <110> APPLICANT: YOUNG, ANDREW
 4 GEDULIN, BRONISLAVA
 6 <120> TITLE OF INVENTION: METHODS FOR GLUCAGON SUPPRESSION
 8 <130> FILE REFERENCE: 030639.0031.UTL (249/167 US)
 10 <140> CURRENT APPLICATION NUMBER: 09/889,331B
 C--> 11 <141> **CURRENT FILING DATE: 2001-12-18**
 13 <150> PRIOR APPLICATION NUMBER: PCT/US00/00942
 14 <151> PRIOR FILING DATE: 2000-01-14
 16 <150> PRIOR APPLICATION NUMBER: 60/116,380
 17 <151> PRIOR FILING DATE: 1999-01-14
 19 <150> PRIOR APPLICATION NUMBER: 60/132,017
 20 <151> PRIOR FILING DATE: 1999-04-30
 22 <150> PRIOR APPLICATION NUMBER: 60/175,365
 23 <151> PRIOR FILING DATE: 2000-01-10
 25 <160> NUMBER OF SEQ ID NOS: 239
 27 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 W--> 28 **Microsoft Word 97**
 30 <210> SEQ ID NO: 1
 31 <211> LENGTH: 39
 32 <212> TYPE: PRT
 33 <213> ORGANISM: Heloderma Horridum
 35 <220> FEATURE:
 36 <221> NAME/KEY: AMIDATION
 37 <222> LOCATION: (39)
 38 <223> OTHER INFORMATION: Ser in position 39 is amidated
 40 <400> SEQUENCE: 1
 41 His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 42 1 5 10 15
 44 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 45 20 25 30
 47 Ser Gly Ala Pro Pro Pro Ser
 48 35
 50 <210> SEQ ID NO: 2
 51 <211> LENGTH: 39
 52 <212> TYPE: PRT
 53 <213> ORGANISM: Heloderma Suspectum
 55 <220> FEATURE:
 56 <221> NAME/KEY: AMIDATION
 57 <222> LOCATION: (39)
 58 <223> OTHER INFORMATION: Ser in position 39 is amidated
 60 <400> SEQUENCE: 2
 61 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 62 1 5 10 15

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64 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
65           20           25           30
67 Ser Gly Ala Pro Pro Pro Ser
68           35
70 <210> SEQ ID NO: 3
71 <211> LENGTH: 30
72 <212> TYPE: PRT
73 <213> ORGANISM: Artificial Sequence
75 <220> FEATURE:
76 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
77     Amino Acid Sequence
79 <400> SEQUENCE: 3
80 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
81 1           5           10           15
83 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
84           20           25           30
86 <210> SEQ ID NO: 4
87 <211> LENGTH: 30
88 <212> TYPE: PRT
89 <213> ORGANISM: Artificial Sequence
91 <220> FEATURE:
92 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
93     Amino Acid Sequence
95 <220> FEATURE:
96 <221> NAME/KEY: AMIDATION
97 <222> LOCATION: (30)
98 <223> OTHER INFORMATION: Gly in position 30 is amidated
100 <400> SEQUENCE: 4
101 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
102 1           5           10           15
104 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
105           20           25           30
107 <210> SEQ ID NO: 5
108 <211> LENGTH: 30
109 <212> TYPE: PRT
110 <213> ORGANISM: Artificial Sequence
112 <220> FEATURE:
113 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
114     Construct
116 <220> FEATURE:
117 <221> NAME/KEY: MOD_RES
118 <222> LOCATION: (30)
119 <223> OTHER INFORMATION: AMIDATION, Position 30 is Gly-NH2
121 <400> SEQUENCE: 5
122 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
123 1           5           10           15
125 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly
126           20           25           30
128 <210> SEQ ID NO: 6

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Input Set : A:\249-167 subst seq listing.txt

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129 <211> LENGTH: 28
130 <212> TYPE: PRT
131 <213> ORGANISM: Artificial Sequence
133 <220> FEATURE:
134 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
135     Construct
137 <220> FEATURE:
138 <221> NAME/KEY: MOD_RES
139 <222> LOCATION: (28)
140 <223> OTHER INFORMATION: AMIDATION, Position 28 is Asn-NH2
142 <400> SEQUENCE: 6
143 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
144 1           5           10           15
146 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn
147           20           25
149 <210> SEQ ID NO: 7
150 <211> LENGTH: 39
151 <212> TYPE: PRT
152 <213> ORGANISM: Artificial Sequence
154 <220> FEATURE:
155 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
156     Construct
158 <220> FEATURE:
159 <221> NAME/KEY: MOD_RES
160 <222> LOCATION: (30)
161 <223> OTHER INFORMATION: AMIDATION, Position 30 is Gly-NH2
163 <400> SEQUENCE: 7
164 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
165 1           5           10           15
167 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
168           20           25           30
170 Ser Gly Ala Pro Pro Pro Ser
171           35
173 <210> SEQ ID NO: 8
174 <211> LENGTH: 28
175 <212> TYPE: PRT
176 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
180     Construct
182 <220> FEATURE:
183 <221> NAME/KEY: MOD_RES
184 <222> LOCATION: (28)
185 <223> OTHER INFORMATION: AMIDATION, Position 28 is Asn-NH2
187 <400> SEQUENCE: 8
188 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
189 1           5           10           15
191 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn
192           20           25

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194 <210> SEQ ID NO: 9
195 <211> LENGTH: 28
196 <212> TYPE: PRT
197 <213> ORGANISM: Artificial Sequence
199 <220> FEATURE:
200 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
201     Construct
203 <220> FEATURE:
204 <221> NAME/KEY: MOD_RES
205 <222> LOCATION: (28)
206 <223> OTHER INFORMATION: AMIDATION, Position 28 is Asn-NH2
208 <400> SEQUENCE: 9
209 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
210 1      5      10      15
212 Ala Val Arg Leu Ala Ile Glu Phe Leu Lys Asn
213      20      25
216 <210> SEQ ID NO: 10
217 <211> LENGTH: 39
218 <212> TYPE: PRT
219 <213> ORGANISM: Artificial Sequence
221 <220> FEATURE:
222 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
223     Construct
225 <220> FEATURE:
226 <221> NAME/KEY: MOD_RES
227 <222> LOCATION: (39)
228 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2
230 <400> SEQUENCE: 10
231 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
232 1      5      10      15
234 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
235      20      25      30
237 Ser Gly Ala Pro Pro Pro Ser
238      35
240 <210> SEQ ID NO: 11
241 <211> LENGTH: 39
242 <212> TYPE: PRT
243 <213> ORGANISM: Artificial Sequence
245 <220> FEATURE:
246 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
247     Construct
249 <220> FEATURE:
250 <221> NAME/KEY: MOD_RES
251 <222> LOCATION: (39)
252 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2
254 <400> SEQUENCE: 11
255 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
256 1      5      10      15
258 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser

```

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259          20          25          30
261 Ser Gly Ala Pro Pro Pro Ser
262          35
264 <210> SEQ ID NO: 12
265 <211> LENGTH: 39
266 <212> TYPE: PRT
267 <213> ORGANISM: Artificial Sequence
269 <220> FEATURE:
270 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
271 Construct
273 <220> FEATURE:
274 <221> NAME/KEY: MOD_RES
275 <222> LOCATION: (39)
276 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2
278 <400> SEQUENCE: 12
279 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
280 1          5          10          15
282 Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
283          20          25          30
285 Ser Gly Ala Pro Pro Pro Ser
286          35
288 <210> SEQ ID NO: 13
289 <211> LENGTH: 39
290 <212> TYPE: PRT
291 <213> ORGANISM: Artificial Sequence
293 <220> FEATURE:
294 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
295 Construct
297 <220> FEATURE:
298 <221> NAME/KEY: MOD_RES
299 <222> LOCATION: (39)
300 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2
302 <400> SEQUENCE: 13
303 Tyr Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
304 1          5          10          15
306 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
307          20          25          30
309 Ser Gly Ala Pro Pro Pro Ser
310          35
312 <210> SEQ ID NO: 14
313 <211> LENGTH: 39
314 <212> TYPE: PRT
315 <213> ORGANISM: Artificial Sequence
317 <220> FEATURE:
318 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
319 Construct
321 <220> FEATURE:
322 <221> NAME/KEY: MOD_RES
323 <222> LOCATION: (39)

```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/889,331B

DATE: 01/14/2005
TIME: 17:11:44

Input Set : A:\249-167 subst seq listing.txt
Output Set: N:\CRF4\01142005\I889331B.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:16; Xaa Pos. 6
Seq#:21; Xaa Pos. 10
Seq#:22; Xaa Pos. 10
Seq#:23; Xaa Pos. 14
Seq#:24; Xaa Pos. 14
Seq#:25; Xaa Pos. 22
Seq#:29; Xaa Pos. 23
Seq#:32; Xaa Pos. 31,36,37,38
Seq#:33; Xaa Pos. 36,37,38
Seq#:34; Xaa Pos. 31,36,37,38
Seq#:35; Xaa Pos. 36,37,38
Seq#:36; Xaa Pos. 31,36,37,38
Seq#:37; Xaa Pos. 31,36,37,38
Seq#:38; Xaa Pos. 31,36,37,38
Seq#:39; Xaa Pos. 36,37,38
Seq#:40; Xaa Pos. 31,36,37,38
Seq#:41; Xaa Pos. 1,2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23
Seq#:41; Xaa Pos. 24,25,26,27,28,31,36,37,38
Seq#:42; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23
Seq#:42; Xaa Pos. 24,25,26,27,28,31,36,37,38,39
Seq#:43; Xaa Pos. 1,2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23,24
Seq#:43; Xaa Pos. 25,26,27,28,31,36,37,38
Seq#:44; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23
Seq#:44; Xaa Pos. 24,25,26,27,28,31,36,37,38
Seq#:45; Xaa Pos. 1,2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,18,20,21,22,23,24
Seq#:45; Xaa Pos. 25,26,27,28,31,36,37,38
Seq#:46; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23
Seq#:46; Xaa Pos. 24,25,26,27,28,31,36,37,38,39
Seq#:47; Xaa Pos. 1,2,3,6,7,8,9,10,14,22,23,24,25,31,36,37,38,39
Seq#:48; Xaa Pos. 1,2,3,6,7,8,9,10,14,22,23,24,25,27,28,31,36,37,38,39
Seq#:91; Xaa Pos. 31,36,37,38
Seq#:92; Xaa Pos. 36,37,38
Seq#:93; Xaa Pos. 31
Seq#:94; Xaa Pos. 31,36,37
Seq#:95; Xaa Pos. 31,36,37
Seq#:96; Xaa Pos. 31,36
Seq#:99; Xaa Pos. 6
Seq#:103; Xaa Pos. 10
Seq#:104; Xaa Pos. 22
Seq#:105; Xaa Pos. 23
Seq#:109; Xaa Pos. 31,36,37
Seq#:110; Xaa Pos. 1,26
Seq#:111; Xaa Pos. 1,26
Seq#:112; Xaa Pos. 1,26

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Seq#:113; Xaa Pos. 1,26
Seq#:114; Xaa Pos. 1,27
Seq#:115; Xaa Pos. 1,27
Seq#:116; Xaa Pos. 1,27
Seq#:117; Xaa Pos. 1,27
Seq#:133; Xaa Pos. 6
Seq#:134; Xaa Pos. 6

VERIFICATION SUMMARY

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L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:28 M:259 W: Allowed number of lines exceeded, <170> SOFTWARE:
L:380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:505 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:534 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:563 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:619 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:16
L:720 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:16
L:802 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:16
M:341 Repeated in SeqNo=32
L:834 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:32
L:865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:16
M:341 Repeated in SeqNo=34
L:897 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:32
L:928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:16
M:341 Repeated in SeqNo=36
L:962 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:16
M:341 Repeated in SeqNo=37
L:996 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:16
M:341 Repeated in SeqNo=38
L:1028 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:32
L:1059 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:16
M:341 Repeated in SeqNo=40
L:1262 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0
M:341 Repeated in SeqNo=41
L:1478 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:0
M:341 Repeated in SeqNo=42
L:1677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0
M:341 Repeated in SeqNo=43
L:1890 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:0
M:341 Repeated in SeqNo=44
L:2103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:0
M:341 Repeated in SeqNo=45
L:2330 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 after pos.:0
M:341 Repeated in SeqNo=46
L:2442 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:0
M:341 Repeated in SeqNo=47
L:2566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:0
M:341 Repeated in SeqNo=48
L:3521 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:91 after pos.:16
M:341 Repeated in SeqNo=91
L:3554 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:92 after pos.:32
L:3580 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93 after pos.:16
L:3614 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:16
M:341 Repeated in SeqNo=94
L:3649 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95 after pos.:16
M:341 Repeated in SeqNo=95

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L:3683 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:96 after pos.:16

M:341 Repeated in SeqNo=96

L:3754 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99 after pos.:0

L:3843 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:103 after pos.:0